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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/730,469

DATE: 12/26/2000
TIME: 13:40:37

ENTERED

Input Set : A:\seqlist.txt
Output Set: N:\CRF3\12262000\I730469.raw

```
 4 <110> APPLICANT: Anthony P. Heaney
 5             Gregory A. Horwitz
 6             Xun Zhang
 7             Shlomo Melmed
 9 <120> TITLE OF INVENTION: Methods of Using Pituitary Tumor
10        Transforming Gene (PTTG) Carboxy-terminal Peptides to
11        Inhibit Neoplastic Cellular Proliferation And/Or
12        Transformation of Breast and Ovarian Cells
15 <130> FILE REFERENCE: CEDAR-45257
C--> 17 <140> CURRENT APPLICATION NUMBER: US/09/730,469
18 <141> CURRENT FILING DATE: 2000-12-04
20 <150> PRIOR APPLICATION NUMBER: US CIP 09/687,911
21 <151> PRIOR FILING DATE: 2000-10-13
23 <150> PRIOR APPLICATION NUMBER: US CIP 09/569,956
24 <151> PRIOR FILING DATE: 2000-05-12
26 <150> PRIOR APPLICATION NUMBER: US 08/894,251
27 <151> PRIOR FILING DATE: 1999-07-23
29 <150> PRIOR APPLICATION NUMBER: PCT/US97/21463
30 <151> PRIOR FILING DATE: 1997-11-21
32 <150> PRIOR APPLICATION NUMBER: US 60/031,338
33 <151> PRIOR FILING DATE: 1996-11-21
35 <160> NUMBER OF SEQ ID NOS: 19
37 <170> SOFTWARE: FastSEQ for Windows Version 4.0
39 <210> SEQ ID NO: 1
40 <211> LENGTH: 974
41 <212> TYPE: DNA
42 <213> ORGANISM: Rattus rattus
44 <400> SEQUENCE: 1
45 aattcggcac gagccaacct tgagcatctg atcctcttgg cttctccttc ctatcgctga 60
46 gctggtaggc tggagacagt tgtttgggtg ccaacatcaa caaacgattt ctgtagttta 120
47 gcgtttatga ccctggcgtg aagatttaag gtctggatta agcctgttga cttctccagc 180
48 tacttctaaa tttttgtgca taggtgctct ggtctctgtt gctgcttagt tcttccagcc 240
49 ttcctcaatg ccagtttat aatatgcagg tctctcccct cagtaatcca ggatggctac 300
50 tctgatcttt gttgataagg ataacgaaga gccaggcagc cgtttggcat ctaaggatgg 360
51 attgaagctg ggctctggtg tcaaagcctt agatgggaaa ttgcaggttt caacgccacg 420
52 agtcggcaaa gtgttcggtg ccccaggctt gcctaaagcc agcaggaagg ctctgggaac 480
53 tgtcaacaga gttactgaaa agccagtgaa gagtagtaaa cccctgcaat cgaaacagcc 540
54 gactctgagt gtgaaaaaga tcaccgagaa gtctactaag acacaaggct ctgctcctgc 600
55 tcctgatgat gcctacccag aaatagaaaa gttcttcgcc ttcgatcctc tagattttga 660
56 gagtttttgac ctgcctgaag agcaccagat ctcacttctc cccttgaatg gagtgcctct 720
57 catgatcctg aatgaagaga gggggcttga gaagctgctg cacctggacc cccttcccc 780
58 tctgcagaag cccttcctac cgtgggaatc tgatccgttg ccgtctcctc ccagcgccct 840
59 ctccgctctg gatgttgaat tgccgcctgt ttgttacgat gcagatattt aaacgtctta 900
60 ctcctttata gtttatgtaa gttgtattaa taaagcattt gtgtgtaaaa aaaaaaaaa 960
61 aaactcgaga gtac                                              974
63 <210> SEQ ID NO: 2
64 <211> LENGTH: 199
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                    RAW SEQUENCE LISTING            DATE: 12/26/2000
                    PATENT APPLICATION:  US/09/730,469    TIME: 13:40:37

                    Input Set : A:\seqlist.txt
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 65  <212> TYPE: PRT
 66  <213> ORGANISM: Rattus rattus
 68  <400> SEQUENCE: 2
 69  Met Ala Thr Leu Ile Phe Val Asp Lys Asp Asn Glu Glu Pro Gly Ser
 70   1               5                  10                  15
 71  Arg Leu Ala Ser Lys Asp Gly Leu Lys Leu Gly Ser Gly Val Lys Ala
 72              20                  25                  30
 73  Leu Asp Gly Lys Leu Gln Val Ser Thr Pro Arg Val Gly Lys Val Phe
 74          35                  40                  45
 75  Gly Ala Pro Gly Leu Pro Lys Ala Ser Arg Lys Ala Leu Gly Thr Val
 76      50                  55                  60
 77  Asn Arg Val Thr Glu Lys Pro Val Lys Ser Ser Lys Pro Leu Gln Ser
 78  65                  70                  75                  80
 79  Lys Gln Pro Thr Leu Ser Val Lys Lys Ile Thr Glu Lys Ser Thr Lys
 80                  85                  90                  95
 81  Thr Gln Gly Ser Ala Pro Ala Pro Asp Asp Ala Tyr Pro Glu Ile Glu
 82                  100                 105                 110
 83  Lys Phe Phe Pro Phe Asp Pro Leu Asp Phe Glu Ser Phe Asp Leu Pro
 84          115                 120                 125
 85  Glu Glu His Gln Ile Ser Leu Leu Pro Leu Asn Gly Val Pro Leu Met
 86      130                 135                 140
 87  Ile Leu Asn Glu Glu Arg Gly Leu Glu Lys Leu Leu His Leu Asp Pro
 88  145                 150                 155                 160
 89  Pro Ser Pro Leu Gln Lys Pro Phe Leu Pro Trp Glu Ser Asp Pro Leu
 90                  165                 170                 175
 91  Pro Ser Pro Pro Ser Ala Leu Ser Ala Leu Asp Val Glu Leu Pro Pro
 92                  180                 185                 190
 93  Val Cys Tyr Asp Ala Asp Ile
 94              195
 97  <210> SEQ ID NO: 3
 98  <211> LENGTH: 779
 99  <212> TYPE: DNA
100  <213> ORGANISM: Homo sapiens
102  <400> SEQUENCE: 3
103  atggccgcga gttgtggttt aaaccaggag tgccgcgcgt ccgttcaccg cggcctcaga  60
104  tgaatgcggc tgttaagacc tgcaataatc cagaatggct actctgatct atgttgataa  120
105  ggaaaatgga gaaccaggca cccgtgtggt tgctaaggat gggctgaagc tggggtctgg  180
106  accttcaatc aaagccttag atgggagatc tcaagtttca acaccacgtt ttggcaaaac  240
107  gttcgatgcc ccaccagcct tacctaaagc tactagaaag gcttgggaa ctgtcaacag  300
108  agctacagaa aagtctgtaa agaccaaggg acccctcaaa caaaaacagc caagctttc  360
109  tgccaaaaag atgactgaga agactgttaa agcaaaaagc tctgttcctg cctcagtga  420
110  tgcctatcca gaaatagaaa aattctttcc cttcaatcct ctagactttg agagttttga  480
111  cctgcctgaa gagcaccaga ttgcgcacct cccttgagt ggagtgcctc tcatgatcct  540
112  tgacgaggag agagagcttg aaaagctgtt tcagctgggc ccccttcac ctgtgaagat  600
113  gccctctcca ccatgggaat ccaatctgtt gcagtctcct tcaagcattc tgtcgaccct  660
114  ggatgttgaa ttgccacctg tttgctgtga catagatatt taaatttctt agtgcttcag  720
115  agtttgtgtg tatttgtatt aataaagcat tctttaacag ataaaaaaaa aaaaaaaaa  779
117  <210> SEQ ID NO: 4
118  <211> LENGTH: 202
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119 <212> TYPE: PRT
120 <213> ORGANISM: Homo sapiens
122 <400> SEQUENCE: 4
123 Met Ala Thr Leu Ile Tyr Val Asp Lys Glu Asn Gly Glu Pro Gly Thr
124  1               5                  10                  15
125 Arg Val Val Ala Lys Asp Gly Leu Lys Leu Gly Ser Gly Pro Ser Ile
126              20                  25                  30
127 Lys Ala Leu Asp Gly Arg Ser Gln Val Ser Thr Pro Arg Phe Gly Lys
128          35                  40                  45
129 Thr Phe Asp Ala Pro Pro Ala Leu Pro Lys Ala Thr Arg Lys Ala Leu
130          50                  55                  60
131 Gly Thr Val Asn Arg Ala Thr Glu Lys Ser Val Lys Thr Lys Gly Pro
132 65                  70                  75                  80
133 Leu Lys Gln Lys Gln Pro Ser Phe Ser Ala Lys Lys Met Thr Glu Lys
134                  85                  90                  95
135 Thr Val Lys Ala Lys Ser Ser Val Pro Ala Ser Asp Asp Ala Tyr Pro
136                 100                 105                 110
137 Glu Ile Glu Lys Phe Phe Pro Phe Asn Pro Leu Asp Phe Glu Ser Phe
138                 115                 120                 125
139 Asp Leu Pro Glu Glu His Gln Ile Ala His Leu Pro Leu Ser Gly Val
140         130                 135                 140
141 Pro Leu Met Ile Leu Asp Glu Glu Arg Glu Leu Glu Lys Leu Phe Gln
142 145                 150                 155                 160
143 Leu Gly Pro Pro Ser Pro Val Lys Met Pro Ser Pro Pro Trp Glu Ser
144                 165                 170                 175
145 Asn Leu Leu Gln Ser Pro Ser Ser Ile Leu Ser Thr Leu Asp Val Glu
146                 180                 185                 190
147 Leu Pro Pro Val Cys Cys Asp Ile Asp Ile
148         195                 200
151 <210> SEQ ID NO: 5
152 <211> LENGTH: 31
153 <212> TYPE: DNA
154 <213> ORGANISM: Artificial Sequence
156 <220> FEATURE:
157 <223> OTHER INFORMATION: Synthetic oligonucleotide.
159 <400> SEQUENCE: 5
160 gatgctctcc gcactctggg aatccaatct g                          31
162 <210> SEQ ID NO: 6
163 <211> LENGTH: 32
164 <212> TYPE: DNA
165 <213> ORGANISM: Artificial Sequence
167 <220> FEATURE:
168 <223> OTHER INFORMATION: Synthetic oligonucleotide.
170 <400> SEQUENCE: 6
171 ttcacaagtt gaggggcgcc cagctgaaac ag                         32
173 <210> SEQ ID NO: 7
174 <211> LENGTH: 32
175 <212> TYPE: DNA
176 <213> ORGANISM: Artificial Sequence
```

```
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```
178 <220> FEATURE:
179 <223> OTHER INFORMATION: Synthetic oligonucleotide specific to pCI-neo
180       plasmid vector.
182 <400> SEQUENCE: 7
183 ggctagagta cttaatacga ctcactatag gc                          32
185 <210> SEQ ID NO: 8
186 <211> LENGTH: 31
187 <212> TYPE: DNA
188 <213> ORGANISM: Homo sapiens
190 <400> SEQUENCE: 8
191 ctatgtcaca gcaaacaggt ggcaattcaa c                           31
193 <210> SEQ ID NO: 9
194 <211> LENGTH: 56
195 <212> TYPE: PRT
196 <213> ORGANISM: Homo sapiens
198 <400> SEQUENCE: 9
199 Met Ile Leu Asp Glu Glu Arg Glu Leu Glu Lys Leu Phe Gln Leu Gly
200  1               5                  10                  15
201 Pro Pro Ser Pro Val Lys Met Pro Ser Pro Pro Trp Glu Ser Asn Leu
202              20                  25                  30
203 Leu Gln Ser Pro Ser Ser Ile Leu Ser Thr Leu Asp Val Glu Leu Pro
204          35                  40                  45
205 Pro Val Cys Cys Asp Ile Asp Ile
206      50                  55
209 <210> SEQ ID NO: 10
210 <211> LENGTH: 168
211 <212> TYPE: DNA
212 <213> ORGANISM: Homo sapiens
214 <400> SEQUENCE: 10
215 atgatccttg acgaggagag agagcttgaa aagctgtttc agctgggccc cccttcacct 60
216 gtgaagatgc cctctccacc atgggaatcc aatctgttgc agtctccttc aagcattctg 120
217 tcgaccctgg atgttgaatt gccacctgtt tgctgtgaca tagatatt            168
219 <210> SEQ ID NO: 11
220 <211> LENGTH: 16
221 <212> TYPE: DNA
222 <213> ORGANISM: Artificial Sequence
224 <220> FEATURE:
225 <223> OTHER INFORMATION: Anchored primer sequence.
227 <400> SEQUENCE: 11
228 aagcttttt tttttg                                             16
230 <210> SEQ ID NO: 12
231 <211> LENGTH: 13
232 <212> TYPE: DNA
233 <213> ORGANISM: Artificial Sequence
235 <220> FEATURE:
236 <223> OTHER INFORMATION: Arbitrary primer sequence.
238 <400> SEQUENCE: 12
239 aagcttgctg ctc                                               13
241 <210> SEQ ID NO: 13
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242 <211> LENGTH: 16
243 <212> TYPE: DNA
244 <213> ORGANISM: Artificial Sequence
246 <220> FEATURE:
247 <223> OTHER INFORMATION: n = a, g, or c; Anchored primer sequence.
249 <400> SEQUENCE: 13
W--> 250 aagctttttt tttttn                                                   16
252 <210> SEQ ID NO: 14
253 <211> LENGTH: 194
254 <212> TYPE: PRT
255 <213> ORGANISM: Mus musculus
257 <400> SEQUENCE: 14
258 Met Ala Thr Leu Ile Phe Val Asp Lys Asp Asn Glu Glu Pro Gly Arg
259  1               5                   10                  15
260 Arg Leu Ala Ser Lys Asp Gly Leu Lys Leu Gly Thr Gly Val Lys Ala
261              20                  25                  30
262 Leu Asp Gly Lys Leu Gln Val Ser Thr Pro Arg Val Gly Lys Val Phe
263          35                  40                  45
264 Asn Ala Pro Ala Val Pro Lys Ala Ser Arg Lys Ala Leu Gly Thr Val
265      50                  55                  60
266 Asn Arg Val Ala Glu Lys Pro Met Lys Thr Gly Lys Pro Leu Gln Pro
267 65                  70                  75                  80
268 Lys Gln Pro Thr Leu Thr Gly Lys Lys Ile Thr Glu Lys Ser Thr Lys
269              85                  90                  95
270 Thr Gln Ser Ser Val Pro Ala Pro Asp Asp Ala Tyr Pro Glu Ile Glu
271              100                 105                 110
272 Lys Phe Phe Pro Phe Asn Pro Leu Asp Phe Asp Leu Pro Glu Glu His
273          115                 120                 125
274 Gln Ile Ser Leu Leu Pro Leu Asn Gly Val Pro Leu Ile Thr Leu Asn
275      130                 135                 140
276 Glu Glu Arg Gly Leu Glu Lys Leu Leu His Leu Gly Pro Pro Ser Pro
277 145                 150                 155                 160
278 Leu Lys Thr Pro Phe Leu Ser Trp Glu Ser Asp Pro Lys Pro Pro Ser
279              165                 170                 175
280 Ala Leu Ser Thr Leu Asp Val Glu Leu Pro Pro Val Cys Tyr Asp Ala
281              180                 185                 190
282 Asp Ile
286 <210> SEQ ID NO: 15
287 <211> LENGTH: 945
288 <212> TYPE: DNA
289 <213> ORGANISM: Mus musculus
291 <400> SEQUENCE: 15
292 tcttgaactt gttatgtagc agqaggccaa atttgagcat cctcttggct tctctttata 60
293 gcagagattg taggctggag acagttttga tgggtgccaa cataaactga tttctgtaag 120
294 agttgagtgt tttatgaccc tggcgtgcag atttaggatc tggattaagc ctgttgactt 180
295 ctccagctac ttataaattt ttqtgcatag gtgccctggg taaagcttgg tctctgttac 240
296 tgcgtagttt ttccagccgt ctcaatgcca atattcaggc tctctccctt agagtaatcc 300
297 agaatggcta ctcttatctt tgttgataag gataatgaag aacccggccg ccgtttggca 360
298 tctaaggatg ggttgaagct gggcactggt gtcaaggcct tagatgggaa attgcaggtt 420
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VERIFICATION SUMMARY DATE: 12/26/2000
PATENT APPLICATION: US/09/730,469 TIME: 13:40:38

Input Set : A:\seqlist.txt
Output Set: N:\CRF3\12262000\I730469.raw

L:17 M:270 C: Current Application Number differs, Replaced Current Application Number
L:250 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:13
L:250 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:13
L:250 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:13